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JC17 Rec'd PCT/PTO 28 MAR 2005

SEQUENCE LISTING

- <110> ONCOTHERAPY SCIENCE, INC.
 JAPAN AS REPRESENTED BY THE PRESIDENT OF THE UNIVERSITY OF TOKYO
- <120> GENES AND POLYPEPTIDES RELATING TO PROSTATE CANCERS
- <130> ONC-A0216P2
- <150> US 60/414,873
- <151> 2002-09-30
- <160> 28
- <170> PatentIn version 3.1
- ⟨210⟩ 1
- 〈211〉 826
- <212> DNA
- <213> Homo sapiens
- <220>
- <221> CDS
- <222> (332)..(634)
- <223>

| | | | | | | | -, | , | | | | | | | | |
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| <400> | 1 | | | | | | | | | | | | | | | |
| aagggg | ctca | tgcag | gaagca | a gt | tece | ggac | ccg | acac | tct | gggt | agga | ga | ccac | țaaa | acc | 60 |
| cggccc | ctca | aagca | agagg" | t ga | cctt | gccc | tca | tcga | ıgag | cgca | caca | ag | acgc | cact | tgt · | 120 |
| aaaagg | atca | cagat | tggag | a ga | catt | ttgc | cac | acga | ıtga | atca | ıcaca | .cc | acat | ctca | atc | 180 |
| cccgag | cttc | agctg | gcagg | a ca | atgo | tgcc | aga | ıggco | tgg | tcct | caga | gc | tcac | gtaa | agc | 240 |
| atctct | ggtg | tgcag | gtatt | t tt | acto | cgtt | , ttt | gaco | eaaa | gaca | ectg | ;aa | catt | cct | gga | 300 |
| gaaaac | agtg | atgt | ggatc | t ta | ntcaa | attt | | let (| | aca t | | lu | | | | 352 |
| aac tt | g ct | c agc | acc | gtg | agc | ccc | aca | gtg | aaa | gca | ctt | ttt | ggc | aaı | g | 400 |
| Asn Le | u Le | u Ser | Thr | Val | Ser | Pro | Thr | Val | Lys | Ala | Leu | Phe | Gly | Lys | S | |
| | 10 | | | | | 15 | | | | | 20 | | | | | |
| act ag | a gt | c tca | ccg | att | ttc | cct | ttc | tct | cct | cga | tct | cct | ttc | ca | g | 448 |
| Thr Ar | g Va | l Ser | Pro | Ile | Phe | Pro | Phe | Ser | Pro | Arg | Ser | Pro | Phe | G1: | n | |
| 25 | ; | | | | 30 | • | | | | 35 | | | | | | |

cct ctt att ccc cgg act cct ggc tca ccc tgg ggc ccc gtg ggt cca 496
Pro Leu Ile Pro Arg Thr Pro Gly Ser Pro Trp Gly Pro Val Gly Pro
40 45 50 55

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| gct | tct | ccc | ttg | gga | cca | ggc | ttt | cca | ata | ggg | ccc | atg | ggg | ccc | ggt | | 544 | |
|------|------|-------|-------------------|------|------|------|-------------------|------|-------------------|------|-----|------|-----|------|------|----|-------------|---|
| Ala | Ser | Pro | Leu | Gly | Pro | Gly | Phe | Pro | Ile | Gly | Pro | Met | Gly | Pro | Gly | | | |
| | | | | 60 | | | | | 65 | | | | | 70 | | | | |
| | | | | | | | | | | | | | | | | | | |
| aaa | cca | gtt | ggg | ccc | aaa | ggc | cca | atg | ttg | ccc | ctt | ggc | ccc | tca | gga | | 592 | , |
| Lys | Pro | Val | Gly | Pro | Lys | Gly | Pro | Met | Leu | Pro | Leu | Gly | Pro | Ser | Gly | | | |
| | | | 75 | | | | | 80 | | | | | 85 | | | | | |
| | | | | | | | | | | | | | | | | | | |
| cca | gtg | gga | ccc | acg | tca | ccc | tta | ttc | ccc | ttc | tgc | ccc | tga | | | | 634 | Ļ |
| Pro | Val | Gly | Pro | Thr | Ser | Pro | Leu | Phe | Pro | Phe | Cys | Pro | | | | | | |
| | | 90 | | | | | 95 | | | | | 100 | | | | | | |
| | | | | | | | | | | | | • | | | | | | |
| ggc | ccag | tct | ctcc [.] | tegg | ag g | cctt | tctc [.] | t cc | catg | ggcc | ctg | caag | ccc | cttg | gggc | ca | 694 | Ļ |
| | | | | | | | | ì | | | | | | | | ٠ | | |
| tgt | tttc | ctg | ggga [.] | tcct | ct t | gagc | cttg | a tc | acct [.] | ttga | tgc | cttt | tgc | ttca | actt | tt | 7 54 | Ļ |
| | | | | | | | | | | | | | | | | | - | |
| cca | tctg | ctc | ctaa | atag | ag a | aaga | gcaa | a ta | aaga | gata | gtt | tgtg | aaa | gata | aaaa | aa | 814 | Ė |
| | | | | | | | | | | | | | | | | | | |
| aaaa | aaaa | aaa : | aa | | | | | | | | | | | | | | 826 | ; |

⟨210⟩ 2

<211> 100

<212> PRT

<213> Homo sapiens

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⟨400⟩ 2

Met Gly Thr Ser Glu Glu Gly Asn Leu Leu Ser Thr Val Ser Pro Thr

Val Lys Ala Leu Phe Gly Lys Thr Arg Val Ser Pro Ile Phe Pro Phe

Ser Pro Arg Ser Pro Phe Gln Pro Leu Ile Pro Arg Thr Pro Gly Ser

Pro Trp Gly Pro Val Gly Pro Ala Ser Pro Leu Gly Pro Gly Phe Pro

Ile Gly Pro Met Gly Pro Gly Lys Pro Val Gly Pro Lys Gly Pro Met

Leu Pro Leu Gly Pro Ser Gly Pro Val Gly Pro Thr Ser Pro Leu Phe

Pro Phe Cys Pro

⟨210⟩ 3

<211> 6805

| 701 | ი\ | T\NTA |
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| <21 | 4/ | DNA |

<213> Homo sapiens

<220>

<221> CDS

<222> (265).. (3195)

<223>

<400> 3

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cgcggccagg gccgaggcag gcctgacccg gggccgggca gcccgcgcga ctttcggaac 120
atggcaaccc gtgtgtgtct catcccagaa agagaagact ttaaccactg tgatgcctga 180
gaatccagtg tgacgtttct ccagatactt catgctgttc acctgtgtcc tcgccgcacc 240
actgccgcac acgactcctg aacc atg ggg gaa aac gag gat gag aag cag 291

1 5

Met Gly Glu Asn Glu Asp Glu Lys Gln

gcc cag gcg ggg cag gtt ttt gag aac ttt gtc cag gca tcc acg tgc 339

Ala Gln Ala Gly Gln Val Phe Glu Asn Phe Val Gln Ala Ser Thr Cys

10 20 25

aaa ggt acc ctc cag gcc ttc aac att ctc aca cga cac ctg gac cta

387

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Lys Gly Thr Leu Gln Ala Phe Asn Ile Leu Thr Arg His Leu Asp Leu
30 35 40

gac cct ctg gac cac aga aac ttt tat tcc aag ctc aag tcc aag gtg

Asp Pro Leu Asp His Arg Asn Phe Tyr Ser Lys Leu Lys Ser Lys Val

45

50

55

acc acc tgg aaa gcc aaa gcc ctg tgg tac aaa ttg gat aag cgt ggt 483

Thr Thr Trp Lys Ala Lys Ala Leu Trp Tyr Lys Leu Asp Lys Arg Gly
60 65 70

tcc cac aaa gag tat aag cga ggg aag tcg tgc acg aac acc aag tgt 531
Ser His Lys Glu Tyr Lys Arg Gly Lys Ser Cys Thr Asn Thr Lys Cys
75 80 85

ctc ata gtt ggg gga gga ccc tgt ggc ttg cgc act gcc att gaa ctt 579

Leu Ile Val Gly Gly Pro Cys Gly Leu Arg Thr Ala Ile Glu Leu

90 95 100 105

gcc tac ctg gga gcc aaa gtg gtc gtg gtg gag aag agg gac tcc ttc 627
Ala Tyr Leu Gly Ala Lys Val Val Val Glu Lys Arg Asp Ser Phe
110 115 120

tcc cgg aac aac gtg cta cac ctc tgg cct ttc acc atc cat gac ctt

Ser Arg Asn Asn Val Leu His Leu Trp Pro Phe Thr Ile His Asp Leu

130

125

135

1011

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| | _ | | | | | | | | | | | | | | | |
|------|-----|------|-----|-----------|------|-----|------|------|-----|-----|-------|------|-----|------|-------|-----|
| cgg | ggc | ctg | gga | gcc | aag | aag | ttc | tat | ggg | aag | ttc | tgt | gct | ggc | tcc | 723 |
| Arg | Gly | Leu | Gly | Ala | Lys | Lys | Phe | Tyr | Gly | Lys | Phe | Cys | Ala | Gly | Ser | |
| | | 140 | | | | | 145 | | | | | 150 | | | | |
| | | | | | | | | | | | | | | | - | |
| atc | gac | cat | atc | agt | att | cgc | caa | cta | cag | ctc | atc | cta | ttc | aag | gtg | 771 |
| Ile | Asp | His | Ile | Ser | Ile | Arg | Gln | Leu | Gln | Leu | Ile | Leu | Phe | Lys | Val | |
| | 155 | | | | | 160 | | | | | 165 | | | | | |
| | | | | | | | | | | | | | | | | |
| gcc | ctg | atg | ctg | gga | gtt | gaa | atc | cat | gtg | aat | gtg | gag | ttc | gtg | aag | 819 |
| Ala | Leu | Met | Leu | Gly | Val | Glu | Ile | His | Val | Asn | Val | Glu | Phe | Val | Lys | |
| 170 | | | | | 175 | | | | | 180 | | | | | 185 | , |
| | | | | | | | | | | | | | | | , | • |
| gtt | cta | gag | cct | cct | gaa | gat | caa | gaa | aat | caa | aaa | att | ggc | tgg | cgg | 867 |
| Val | Leu | Glu | Pro | Pro | Glu | Asp | Gln | Glu | Asn | Gln | Lys | Ile | Gly | Trp | Arg | |
| | | | | 190 | | | | | 195 | | | | | 200 | | |
| | | , | | | | | | | | | | | * | | . • | |
| gca | gaa | ttt | ctc | cct | aca | gac | cat | tct | ctg | tcg | gag | ttt | gag | ttt | gac | 915 |
| | Glu | | | | | , | | | | | | | | | | |
| **** | ulu | 1110 | 205 | 110 | | пор | 1110 | 210 | Dog | 501 | 014 | | 215 | 11.0 | IIO D | |
| | | | 200 | | | | | 210 | | | | | 210 | | | |
| at a | ota | 0++ | aat | ~~ | ant. | aao | 000 | aaa | 200 | 200 | o t a | go o | aaa | ++0 | 0.00 | 963 |
| • | atc | | | | | | | | | | | | | | | 903 |
| Val | Ile | | ΩŢΆ | RIA | ASP | ΩŢΆ | | vi.Ř | ASN | inr | Leu | | ΩŢλ | rne | VLA | |
| | | 220 | | | | | 225 | | | • | | 230 | | | | |

aga aaa gaa ttc cgt ggg aag ctg gcg att gcc atc acc gcc aac ttc

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Arg Lys Glu Phe Arg Gly Lys Leu Ala Ile Ala Ile Thr Ala Asn Phe
235 240 245

ata aac aga aac agc aca gcg gaa gcc aag gtg gaa gag att agt ggt 1059

Ile Asn Arg Asn Ser Thr Ala Glu Ala Lys Val Glu Glu Ile Ser Gly

250 265 260 265

gtg gct ttc atc ttc aat cag aaa ttt ttt cag gac ctt aaa gaa gaa 1107
Val Ala Phe Ile Phe Asn Gln Lys Phe Phe Gln Asp Leu Lys Glu Glu
270 275 280

aca ggc ata gat ctt gag aac att gtt tac tac aag gac tgc acc cac 1155

Thr Gly Ile Asp Leu Glu Asn Ile Val Tyr Tyr Lys Asp Cys Thr His

285 290 295

tat ttt gta atg aca gcc aag aag cag agc ctg ctc gac aaa ggt gtc 1203

Tyr Phe Val Met Thr Ala Lys Lys Gln Ser Leu Leu Asp Lys Gly Val

300 305 310

atc att aac gac tac atc gac aca gag atg ctg ctg tgt gcg gag aac 1251

Ile Ile Asn Asp Tyr Ile Asp Thr Glu Met Leu Leu Cys Ala Glu Asn

315 320 325

gtg aac caa gac aac ctg cta tcc tat gcc cgg gaa gct gca gac ttt 1299
Val Asn Gln Asp Asn Leu Leu Ser Tyr Ala Arg Glu Ala Ala Asp Phe
330 335 340 345

1635

9/59

| gcc | acc | aac | tac | cag | ctg | cca | tcc | tta | gac | ttt | gcc | atg | aac | cac | tat | 1347 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| Ala | Thr | Asn | Tyr | Gln | Leu | Pro | Ser | Leu | Asp | Phe | Ala | Met | Asn | His | Tyr | |
| | | | | 350 | | | | | 355 | | | | | 360 | | |
| | | | | | | | | | | | | | | | | |
| ggg | cag | cct | gat | gtg | gcc | atg | ttt | gac | ttt | acc | tgc | atg | tat | gcc | tca | 1395 |
| Gly | Gln | Pro | Asp | Val | Ala | Met | Phe | Asp | Phe | Thr | Cys | Met | Tyr | Ala | Ser | |
| | | | 365 | • | | | | 370 | | | | | 375 | | | |
| | | | | | | | | | | | | | | | | |
| gag | aac | gcg | gcc | ctg | gtg | cgg | gag | cgg | cag | gcg | cac | cag | ctg | ctc | gtg | 1443 |
| Glu | Asn | Ala | Ala | Leu | Val | Arg | Glu | Arg | Gln | Ala | His | G1n | Leu | Leu | Val | |
| | | 380 | | | | | 385 | | | | | 390 | | | | |
| | ٠. | | | | | | , | | | | | | | | | |
| gcc | ctt | gtg | ggt | gac | agc | ttg | ctt | gag | cca | ttt | tgg | ссс | atg | ggt | aca | 1491 |
| Ala | Leu | Val | Gly | Asp | Ser | Leu | Leu | Glu | Pro | Phe | Trp | Pro | Met | Gly | Thr | |
| | 395 | | | | | 400 | | | | | 405 | | | | | • |
| | | | | | | | | | | | | | | | • | |
| ggc | tgt | gcc | cgt | ggc | ttc | ctg | gca | gcc | ttt | gac | acg | gca | tgg | atg | gtg | 1539 |
| G1y | Cys | Ala | Arg | Gly | Phe | Leu | Ala | Ala | Phe | Asp | Thr | Ala | Trp | Met | Val | |
| 410 | | | | | 415 | | | | | 420 | | | | | 425 | |
| | | | | | | | | | | | | | | | | |
| aag | agc | tgg | aac | cag | ggc | acc | cct | ccc | ctg | gag | ctg | ctg | gct | gaa | agg | 1587 |
| Lys | Ser | Trp | Asn | G1n | Gly | Thr | Pro | Pro | Leu | Glu | Leu | Leu | Ala | Glu | Arg | |
| | | | | 430 | | | | | 435 | | | | | 440 | | |
| | | | | | | | | | | | | | | | | |

gaa agt ctc tac cgg ctg tta cct cag aca acc ccg gag aac atc aac

10/59

Glu Ser Leu Tyr Arg Leu Leu Pro Gln Thr Thr Pro Glu Asn Ile Asn
445
450
455

aag aac ttt gag cag tac acg ttg gac cca ggg aca cgg tac cca aac 1683

Lys Asn Phe Glu Gln Tyr Thr Leu Asp Pro Gly Thr Arg Tyr Pro Asn

460 465 470

ctc aac tca cac tgt gtc agg ccc cat cag gtg aag cat ttg tat atc 1731

Leu Asn Ser His Cys Val Arg Pro His Gln Val Lys His Leu Tyr Ile

475 480 485

act aag gag ctg gag cac tac cct ctc gag aga ctg ggc tcg gtg agg 1779

Thr Lys Glu Leu Glu His Tyr Pro Leu Glu Arg Leu Gly Ser Val Arg

490 495 500 505

aga tct gtc aac ctc tcc agg aag gag tca gat atc cgg ccc agc aag 1827

Arg Ser Val Asn Leu Ser Arg Lys Glu Ser Asp Ile Arg Pro Ser Lys

510 515 520

ctc ctg acc tgg tgc cag cag cag aca gag ggc tac cag cat gtc aac

1875

Leu Leu Thr Trp Cys Gln Gln Gln Thr Glu Gly Tyr Gln His Val Asn

525

530

535

gtc acc gac ctg acc aca tcc tgg cgc agt ggg ttg gcc ctg tgt gcc 1923

Val Thr Asp Leu Thr Thr Ser Trp Arg Ser Gly Leu Ala Leu Cys Ala

540

545

550

2259

11/59

| atc | atc | cac | cgc | ttc | cgg | cct | gag | ctc | atc | aac | ttt | gac | tct | ttg | aat | 1971 |
|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| [le | Ile | His | Arg | Phe | Arg | Pro | Glu | Leu | Ile | Asn | Phe | Asp | Ser | Leu | Asn | |
| | 555 | | | | | 560 | | • | | | 565 | | | | | |
| | | | | | | | | | | | | | | | | |
| gaa | gat | gat | gct | gtg | gag | aac | aac | cag | ctc | gca | ttt | gat | gtg | gcc | gag | 2019 |
| Glu | Asp | Asp | Ala | Val | Glu | Asn | Asn | Gln | Leu | Ala | Phe | Asp | Val | Ala | Glu | |
| 570 | | | | | 575 | | | | | 580 | | | | | 585 | |
| | | | | | | | | | | | | | | | | |
| cga | gag | ttt | ggg | atc | cct | cca | gtg | acc | acg | ggc | aaa | gag | atg | gca | tct | 2067 |
| Arg | Glu | Phe | Gly | Ile | Pro | Pro | Val | Thr | Thr | Gly | Lys | Glu | Met | Ala | Ser | |
| | | | | 590 | | | | | 595 | | | | | 600 | | |
| | | | | | | | • | | | | | | | | | |
| gcc | cag | gag | cct | gac | aag | ctc | agc | atg | gtc | atg | tac | ctc | tcc | aag | ttc | 2115 |
| Ala | Gln | Glu | Pro | Asp | Lys | Leu | Ser | Met | Val | Met | Tyr | Leu | Ser | Lys | Phe | |
| | | | 605 | | | | | 610 | | | | | 615 | | | |
| | | | | ٠ | | | • | | | | | | | | r | |
| tac | gag | ctc | ttc | cgg | ggc | acc | cca | ctg | agg | ccc | gtg | gat | tct | tgg | cgc | 2163 |
| Гу́г | Glu | Leu | Phe | Arg | Gly | Thr | Pro | Leu | Arg | Pro | Val | Asp | Ser | Trp | Arg | |
| | | 620 | | | | | 625 | | | | | 630 | | | ٠ | |
| | • | | | | | | | | | | | | | | | |
| aaa | aac | tat | gga | gaa | aat | gct | gac | ctc | agc | ttg | gcc | aaa | tca | tcc | att | 2211 |
| Lys | Asn | Tyr | Gly | Glu | Asn | Ala | Asp | Leu | Ser | Leu | Ala | Lys | Ser | Ser | Ile | |
| | 635 | | | | | 640 | | | | | 645 | | | | | |
| | | | | | | | | | | | | | | | | |

tct aat aac tat ctc aac ctc aca ttt cca agg aag agg act cca cgg

Ser Asn Asn Tyr Leu Asn Leu Thr Phe Pro Arg Lys Arg Thr Pro Arg gtg gat ggt caa acc gga gag aat gac atg aac aaa cgg aga cgg aag Val Asp Gly Gln Thr Gly Glu Asn Asp Met Asn Lys Arg Arg Arg Lys ggc ttc acc aac ctg gac gag cct tca aac ttt tcc agc cgt agc ttg Gly Phe Thr Asn Leu Asp Glu Pro Ser Asn Phe Ser Ser Arg Ser Leu ggc tcc aat caa gag tgt ggg agc agt aag gaa ggt gga aat cag aac Gly Ser Asn Gln Glu Cys Gly Ser Ser Lys Glu Gly Gly Asn Gln Asn aaa gtc aag tcc atg gcg aat cag ctg ctg gcc aag ttt gag gag agc Lys Val Lys Ser Met Ala Asn Gln Leu Leu Ala Lys Phe Glu Glu Ser act cgg aac ccc tca ctc atg aag cag gaa aag aag tca cct tca ggg Thr Arg Asn Pro Ser Leu Met Lys Gln Glu Lys Lys Ser Pro Ser Gly

Phe His Phe His Pro Ser His Leu Arg Thr Val His Pro Gln Glu Ser
750 755 760

ttc cat ttt cat ccc agc cat ttg aga aca gtg cat cct cag gaa tct

2883

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| atg | cga | aag | tca | ttt | ccc | ctt | aac | ctg | gga | ggc | agc | gac | acg | tgt | tac | 25 | 95 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----------------|-----|-----|----|----|
| Met | Arg | Lys | Ser | Phe | Pro | Leu | Asn | Leu | Gly | Gly | Ser | Asp | Thr | Cys | Tyr | | |
| | | | 765 | | | | | 770 | | | | | 775 | | | | |
| | | | | | | | | | | | | | | | | | |
| ttc | tgt | aag | aaa | cgt | gtg | tac | gtg | atg | gaa | cgg | ctg | agc | gcc | gag | ggc | 26 | 43 |
| Phe | Cys | Lys | Lys | Arg | Val | Tyr | Val | Met | Glu | Arg | Leu | Ser | Ala | Glu | Gly | | |
| | | 780 | | | | | 785 | | | | | 790 | | | | | |
| | | | | | | | | | | | | | | | | | |
| cac | ttc | ttc | cac | cgg | gag | tgt | ttc | cgc | tgc | agc | atc | tgt | gcc | acc | acc | 26 | 91 |
| His | Phe | Phe | His | Arg | Glu | Cys | Phe | Arg | Cys | Ser | Ile | Cys | Ala | Thr | Thr | | |
| | 795 | | | | | 800 | | | | | 805 | | | | | | |
| | | | | | | | | | | | | | , | | | | |
| ttg | cgc | ctg | gcc | gcc | tac | acc | ttt | gac | tgc | gat | gaa | ggc | aaa | ttt | tac | 27 | 39 |
| Leu | Arg | Leu | Ala | Ala | Tyr | Thr | Phe | Asp | Cys | Asp | Glu | Gly | Lys | Phe | Tyr | | |
| 810 | | | | | 815 | | | | | 820 | | | | | 825 | | |
| | | | | | | | ` | | | | | | | | | | |
| tgc | aag | cct | cac | ttc | att | cac | tgt | aaa | acc | aat | agc | aaa | caa | cgg | aag | 27 | 87 |
| Cys | Lys | Pro | His | Phe | Ile | His | Cys | Lys | Thr | Asn | Ser | Lys | Gln | Arg | Lys | | |
| | | | | 830 | | | | | 835 | | | | | 840 | | | |
| | | | | , | | | | | | • | | | | | | | |
| aga | cgg | gca | gag | | aag | caa | caa | aga | gag | gag | gag | gca | aca | tgg | caa | 28 | 35 |
| Arg | Arg | Ala | Glu | Leu | Lys | Gln | Gln | Arg | Glu | Glu | Glu | Ala | Thr | Trp | Gln | | |
| | | | 845 | | | | | 850 | | | | | 85 ⁵ | | | | |
| | | | | | • | | | | | | | | | | | | |
| | | | | | | | | | | | | | | | | | |

gag cag gaa gcc cct cgg aga gac act ccc acc gaa agt tct tgc gca

| Glu Gln | Glu Ala | Pro A | rg Arg | Asp | Thr | Pro | Thr | Glu | Ser | Ser | Cys | Ala |
|---------|---------|-------|--------|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | 860 | | | 865 | | | | | 870 | | | |

| gtg | gcc | gcc | att | ggc | acc | ctg | gaa | ggc | agc | ccc | cca | ggt | atc | tcc | acc | 2 | 931 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|---|-----|
| Val | Ala | Ala | Ile | Gly | Thr | Leu | Glu | Gly | Ser | Pro | Pro | Ģly | Ile | Ser | Thr | | |
| | 875 | | | | | 880 | | | | | 885 | | | | | | |

| tcc ttc ttt | agg aag gtg | ctg ggc tgg | ccc ctc agg ctg | ccg agg gac 2979 |
|-------------|-------------|-------------|-----------------|------------------|
| Ser Phe Phe | Arg Lys Val | Leu Gly Trp | Pro Leu Arg Leu | Pro Arg Asp |
| 890 | 895 | | 900 | 905 |

| ctg | tgt | aac | tgg | atg | cag | gga | ctc | ctg | caa | gct | gct | ggc | ctc | cat | atc | 3027 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| Leu | Cys | Asn | Trp | Met | Gln | Gly | Leu | Leu | Gln | Ala | Ala | Gly | Leu | His | Ile | |
| | | | | 910 | | | | | 915 | | | | | 920 | | |

| agg | gac | aat | gct | tac | aac | tac | tgc | tac | atg | tac | gag | ctc | ctg | agc | ctg | 3075 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| Arg | Asp | Asn | Ala | Tyr | Asn | Tyr | Cys | Tyr | Met | Tyr | G1u | Leu | Leu | Ser | Leu | |
| | | | 925 | | | | | 930 | | | | | 935 | | | |

| ggg | ctg | cca | ctc | ctc | tgg | gcg | ttc | tct | gag | gtc | ctg | gca | gcc | atg | tac | 3123 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| Gly | Leu | Pro | Leu | Leu | Trp | Ala | Phe | Ser | Glu | Val | Leu | Ala | Ala | Met | Tyr | |
| • | | 940 | | | | | 945 | | | - | | 950 | | | | |

| agg | gaa | tct | gag | ggc | tcc | ctc | gag | agc | atc | tgc | aac | tgg | gtg | ctc | agg | 3171 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|
| Arg | Glu | Ser | Glu | Gly | Ser | Leu | Glu | Ser | Ile | Cys | Asn | Trp | Val | Leu | Arg | |
| | 955 | | | | | 960 | | | | | 965 | | | | | |

| tgc ttc cca gtc aag ctc | cgc tga c | atggctggc t | gccccaaag | tgccttcaca | 3225 |
|-------------------------|------------|-------------|------------|-------------|------|
| Cys Phe Pro Val Lys Leu | ı Arg | • | | | |
| 970 975 | 5 | | | | |
| | | | | | |
| tttccaggga ggcttcagat g | ggcagtgcgt | ttgcagtttg | ctcaggctct | ggccaggaag | 3285 |
| cctagcattc tctaagcaat t | tagctcaaag | ccaaagaatt | tcacatgggc | cacctccgcc | 3345 |
| tggccttatc agggtgaaca t | tctactcacg | gtgctagggc | cagggatgat | atgaaggatc | 3405 |
| ttttctatag ctttgtgagc c | catacttctg | ggtttacatt | tcaatttttt | taattttaat | 3465 |
| tagcccagag aaagcatttt t | tttctatgag | tgtcaatttt | tctaaacatg | ggtttgaagc | 3525 |
| ttataaccag ttttataaac c | eccttgaaca | ctgcagtgag | ttatcaaagc | .cactgcctgc | 3585 |
| aaagtggatg atttaagatt t | ttacacgcat | gaaaatgagt | gtgccatctc | ctgaccagtg | 3645 |
| ccttttgact taggtaccca g | gatgecaett | gtcagcagca | ggatactttt | tacaacacga | 3705 |
| aagcataatt attttagaag a | aagagagtag | aagggcagaa | tagaattcaa | cttacagaag | 3765 |
| cacggagcag tgtgtggttg g | gctgttatct | gtcccctgg | gaggaggact | gttttgctcc | 3825 |
| cttgttttga tgttaaacag t | tagcttaaag | gctttcccc | ccataccaac | tcacagccaa | 3885 |

atgacaaaga accgtggggt ttcaacagat tctacaaaca tgcattttcc cttcccacta 3945 4005 atgggcactg cagggaaagc ccattggcat ttgaccatgg agctgatgca gtgccaaaga tgagetettt caactgatgg cattttagee eetgtggete ceageggate eeceageeeg 4065 4125 ggctgcaggc tgagccaagg ctgtgcaggg tccatattgg tcaggccaag tggagtggaa 4185 gactetgtee acttatgtgg tgteetttgg gactgagggg gtttgttage acateagget 4245 attgctggga agcgtggcct gcccagtgag cattgcctgt ggacatcctg actgcttagc 4305 tgctccgctg ccacacatat gtggtcaaaa cagaaaccaa tttcacactg ccctgggaaa 4365 ggaatgggtc tgacctccag gggaagctct accatatctt gactggcagg gaaggctggg 4425 agtggaagct atttatggac tgatccaaag gacatatgca tgagtaaggg taaaaatgag 4485 catgcaggtc cacctgtgtt cttactctgg gtatctagaa gagtcctcag ctctccctac 4545 tecaegetge etagacatae acagetgeag ggtetggetg aacaateaag gggeegeeag 4605 agaaaggcca tctacggtgc gcagtgtatc tggagttgct gggcccaaga tagctctgtg 4665 gagttatcac tagagatgcc tctggattaa ctaagaggtg tgcctgggtg tgggtgagga

| gtcagaacct | ttgagagctt | tgagatgaca | gtttctatgg | ggcgggaaga | aggaggtgca | 4725 |
|------------|------------|------------|------------|------------|------------|------|
| tttctacaaa | cacttccctg | aaatccttgg | gaaaaacaga | ggcatggccg | tggccaactc | 4785 |
| tgtgggaact | ggcgcctctg | tccttgttgg | cactgttctc | agtccgatga | cttgcattgt | 4845 |
| gttttctcca | atttttgctg | ggattttaat | gttcagcatg | gtgggaggaa | cccttgattc | 4905 |
| cttttgtttg | agtatagaaa | gtaaattttt | gaggtcatga | tgtgaacggc | catgttattg | 4965 |
| tgattatctt | cagctcagga | taggctgaga | tgctttgtgg | agtgttccat | gaagcccgag | 5025 |
| toggaatoto | tgactgtcgt | gtacagccat | aaggagactg | gtitgaatta | ctgtggcgag | 5085 |
| acagggcgtg | cctgtcagaa | atctgagatg | tttgtacgct | ctgagatgtt | gaacctttct | 5145 |
| ggtgggcagc | accgacaccc | aggggtggac | ccccgaggat | gaatgcctct | aggeeteege | 5205 |
| aacatattca | agaatgaatg | ggagacgcta | gagtaaaatg | ggggcagaga | ggatatcagg | 5265 |
| gagcaagatg | caaactgtgt | gcatccactc | tcgtaaacaa | gtagctggtc | acaaccagaa | 5325 |
| aggticatet | ctcctaagca | aacagcgact | ctttcagagg | aagtttccct | ctttcaatcg | 5385 |
| tggccttatt | ttcaactccg | gtgccttctc | gtgatgttaa | tcatttcctt | ttttccccac | 5445 |

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| а | ctaagctct | cttttctatc | tttctctctc | tttccaatct | tacgccatgg | ccatcagttc | 5505 |
|---|------------|------------|------------|------------|------------|------------|------|
| а | tttcagcct | tccagtgcta | cacccacttc | ttggctgaca | cacttctgct | ctaaggtgac | 5565 |
| t | ggttttctt | gccaattttc | aaagagtggt | actaaccccc | aacccgcttt | ccgcaccccg | 5625 |
| t | cctctccgc | cagcagtact | ggttgcacta | actgtgagtg | tcttgcatac | tgatggactc | 5685 |
| а | tttggtggc | atggttggct | aacagcatgg | cggggggtgt | tcagcttgag | acccatgcct | 5745 |
| g | tgttcattt | cccatggagc | tggcagcctg | gtctacccca | agtgcatgcc | ccgcctctcc | 5805 |
| t | ctctccctt | gggtctgcct | gcgtgcatgc | ttctccagtt | gcgtctgcga | agctacctac | 5865 |
| t | ttcttggga | gggtcgacct | tgatcatgaa | acaataccat | gagggggcct | ctgtcacctt | 5925 |
| t | gaaaagaac | actttttgag | cagcctcaaa | aagctcatac | ataccagcgc | cttcttaaat | 5985 |
| 1 | ggctctaat | gtaaagattg | ttaatgtcat | ttatcaaaac | cataggtgat | tatttggagg | 6045 |
| 8 | gatttaaaaa | acttaattac | tctcaggcct | catcccaagc | ttgacacatg | ctctgtaggt | 6105 |
| 1 | tgaacacata | atcacaaata | ttctagcaaa | tgctgccttg | gttgcagcct | gcactgtaga | 6165 |
| (| ccaagggtt | ttgctgtggc | tcttcttatc | tcccttggct | cataaagccc | cagatgatgc | 6225 |

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| cagagettea | attagagcca | tcatcatccc | aggcagggat | atctttgaga | aatgactcag | 6285 |
|------------|------------|------------|------------|------------|------------|------|
| ttcagcccca | ggcccctgtg | actctgctta | aagcacacat | ttctgctgac | tcttgtacct | 6345 |
| ggggcagcag | gataatcacc | aacacactct | taacgagaaa | caacacacca | agcaccgtgg | 6405 |
| agctgtccta | ggcaacactc | gcggtctcag | gctgcggtgg | gcgtctgtcc | tgcatgtggc | 6465 |
| ccagaccacc | ctgacccccg | ggcctgcctg | cctggccctg | catgctgcac | gctcactgta | 6525 |
| tttgtgcaga | tcctggccag | tacaaagtcg | ttgctcttgt | cttatcttct | cttacagagt | 6585 |
| ctccctccct | ttatagaatg | tcaaccaaag | agtgccctcc | tcccctctca | gcctcctctt | 6645 |
| tagctagcct | ccccatctca | tcacaacgca | tgtctgtgac | ctttggtaat | catttacagt | 6705 |
| gccacacgga | accctgtatt | ttgcacacag | caaaacaaac | aatgtttagc | tttatttatg | 6765 |
| gtatttgatg | ctgtaaatgg | aaataaatat | tgttctttat | | | 6805 |

<210> 4

<211> 976

<212> PRT

<213> Homo sapiens

| 1 | ın | Λ | ` | |
|----|----|---|---|---|
| ٧. | 10 | u | _ | 4 |

Met Gly Glu Asn Glu Asp Glu Lys Gln Ala Gln Ala Gly Gln Val Phe

1 5 10 15

Glu Asn Phe Val Gln Ala Ser Thr Cys Lys Gly Thr Leu Gln Ala Phe
20 25 30

Asn Ile Leu Thr Arg His Leu Asp Leu Asp Pro Leu Asp His Arg Asn

35

40

45

Phe Tyr Ser Lys Leu Lys Ser Lys Val Thr Thr Trp Lys Ala Lys Ala
50 55 60

Leu Trp Tyr Lys Leu Asp Lys Arg Gly Ser His Lys Glu Tyr Lys Arg

75 80

Gly Lys Ser Cys Thr Asn Thr Lys Cys Leu Ile Val Gly Gly Pro

85 90 95

Cys Gly Leu Arg Thr Ala Ile Glu Leu Ala Tyr Leu Gly Ala Lys Val

100 105 110

Val Val Val Glu Lys Arg Asp Ser Phe Ser Arg Asn Asn Val Leu His

115 120 125

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Leu Trp Pro Phe Thr Ile His Asp Leu Arg Gly Leu Gly Ala Lys Lys

130 135 140

Phe Tyr Gly Lys Phe Cys Ala Gly Ser Ile Asp His Ile Ser Ile Arg 145 150 155 160

Gln Leu Gln Leu Ile Leu Phe Lys Val Ala Leu Met Leu Gly Val Glu 165 170 175

Ile His Val Asn Val Glu Phe Val Lys Val Leu Glu Pro Pro Glu Asp
180 185 190

Gln Glu Asn Gln Lys Ile Gly Trp Arg Ala Glu Phe Leu Pro Thr Asp 195 200 205

His Ser Leu Ser Glu Phe Glu Phe Asp Val Ile Ile Gly Ala Asp Gly
210 215 220

Arg Arg Asn Thr Leu Glu Gly Phe Arg Arg Lys Glu Phe Arg Gly Lys
225 230 235 240

Leu Ala Ile Ala Ile Thr Ala Asn Phe Ile Asn Arg Asn Ser Thr Ala
245 250 255

Glu Ala Lys Val Glu Glu Ile Ser Gly Val Ala Phe Ile Phe Asn Gln
260 265 270

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Lys Phe Phe Gln Asp Leu Lys Glu Glu Thr Gly Ile Asp Leu Glu Asn 275 280 285

Ile Val Tyr Tyr Lys Asp Cys Thr His Tyr Phe Val Met Thr Ala Lys
290 295 300

Lys Gln Ser Leu Leu Asp Lys Gly Val IIe IIe Asn Asp Tyr IIe Asp 305 310 315 320

Thr Glu Met Leu Leu Cys Ala Glu Asn Val Asn Gln Asp Asn Leu Leu
325 330 335

Ser Tyr Ala Arg Glu Ala Ala Asp Phe Ala Thr Asn Tyr Gln Leu Pro 340 345 350

Ser Leu Asp Phe Ala Met Asn His Tyr Gly Gln Pro Asp Val Ala Met 355 360 365

Phe Asp Phe Thr Cys Met Tyr Ala Ser Glu Asn Ala Ala Leu Val Arg 370 375 380

Glu Arg Gln Ala His Gln Leu Leu Val Ala Leu Val Gly Asp Ser Leu 385 390 395 400

Leu Glu Pro Phe Trp Pro Met Gly Thr Gly Cys Ala Arg Gly Phe Leu

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405

410

415

Ala Ala Phe Asp Thr Ala Trp Met Val Lys Ser Trp Asn Gln Gly Thr
420 425 430

Pro Pro Leu Glu Leu Leu Ala Glu Arg Glu Ser Leu Tyr Arg Leu Leu
435 440 445

Pro Gln Thr Thr Pro Glu Asn Ile Asn Lys Asn Phe Glu Gln Tyr Thr
450 455 460

Leu Asp Pro Gly Thr Arg Tyr Pro Asn Leu Asn Ser His Cys Val Arg
465 470 475 480

Pro His Gln Val Lys His Leu Tyr Ile Thr Lys Glu Leu Glu His Tyr
485 490 495

Pro Leu Glu Arg Leu Gly Ser Val Arg Arg Ser Val Asn Leu Ser Arg
500 505 510

Lys Glu Ser Asp Ile Arg Pro Ser Lys Leu Leu Thr Trp Cys Gln Gln
515 520 525

Gln Thr Glu Gly Tyr Gln His Val Asn Val Thr Asp Leu Thr Thr Ser
530 535 540

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Trp Arg Ser Gly Leu Ala Leu Cys Ala Ile Ile His Arg Phe Arg Pro 545 550 555 560

Glu Leu Ile Asn Phe Asp Ser Leu Asn Glu Asp Asp Ala Val Glu Asn
565 570 575

Asn Gln Leu Ala Phe Asp Val Ala Glu Arg Glu Phe Gly Ile Pro Pro
580 585 590

Val Thr Thr Gly Lys Glu Met Ala Ser Ala Gln Glu Pro Asp Lys Leu
595 600 605

Ser Met Val Met Tyr Leu Ser Lys Phe Tyr Glu Leu Phe Arg Gly Thr
610 615 620

Pro Leu Arg Pro Val Asp Ser Trp Arg Lys Asn Tyr Gly Glu Asn Ala 625 630 635 640

Asp Leu Ser Leu Ala Lys Ser Ser Ile Ser Asn Asn Tyr Leu Asn Leu
645 650 655

Thr Phe Pro Arg Lys Arg Thr Pro Arg Val Asp Gly Gln Thr Gly Glu
660 665 670

Asn Asp Met Asn Lys Arg Arg Lys Gly Phe Thr Asn Leu Asp Glu 675 680 685

Pro Ser Asn Phe Ser Ser Arg Ser Leu Gly Ser Asn Gln Glu Cys Gly
690 695 700

Ser Ser Lys Glu Gly Gly Asn Gln Asn Lys Val Lys Ser Met Ala Asn 705 710 715 720

Gln Leu Leu Ala Lys Phe Glu Glu Ser Thr Arg Asn Pro Ser Leu Met
725 730 735

Lys Gln Glu Lys Lys Ser Pro Ser Gly Phe His Phe His Pro Ser His
740 745 750

Leu Arg Thr Val His Pro Gln Glu Ser Met Arg Lys Ser Phe Pro Leu
755 760 765

Asn Leu Gly Gly Ser Asp Thr Cys Tyr Phe Cys Lys Lys Arg Val Tyr
770 780

Val Met Glu Arg Leu Ser Ala Glu Gly His Phe Phe His Arg Glu Cys
785 790 795 800

Phe Arg Cys Ser Ile Cys Ala Thr Thr Leu Arg Leu Ala Ala Tyr Thr

805 810 815

Phe Asp Cys Asp Glu Gly Lys Phe Tyr Cys Lys Pro His Phe Ile His

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820

825

830

Cys Lys Thr Asn Ser Lys Gln Arg Lys Arg Arg Ala Glu Leu Lys Gln 835 840 845

Gln Arg Glu Glu Glu Ala Thr Trp Gln Glu Gln Glu Ala Pro Arg Arg 850 855 860

Asp Thr Pro Thr Glu Ser Ser Cys Ala Val Ala Ala Ile Gly Thr Leu 865 870 875 880

Glu Gly Ser Pro Pro Gly Ile Ser Thr Ser Phe Phe Arg Lys Val Leu 885 890 895

Gly Trp Pro Leu Arg Leu Pro Arg Asp Leu Cys Asn Trp Met Gln Gly
900 905 910

Leu Leu Gln Ala Ala Gly Leu His Ile Arg Asp Asn Ala Tyr Asn Tyr
915 920 925

Cys Tyr Met Tyr Glu Leu Leu Ser Leu Gly Leu Pro Leu Leu Trp Ala 930 935 940

Phe Ser Glu Val Leu Ala Ala Met Tyr Arg Glu Ser Glu Gly Ser Leu 945 950 955 960

Glu Ser Ile Cys Asn Trp Val Leu Arg Cys Phe Pro Val Lys Leu Arg
965 970 975

<210> 5

<211> 6742

<212> DNA

<213> Homo sapiens

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<400> 5

cegggeegee tegetegete ceagetetgt cagtggeeeg eggggeeega tegetgegee 60

120

cgcggccagg gccgaggcag gcctgacccg gggccgggca gcccgcgcga ctttcggaac

atggcaaccc gtgtgtgtct catcccagaa agagaagact ttaaccactg tgatgcctga 180

gaatccagtg tgacgtttct ccagatactt catgctgttc acctgtgtcc tcgccgcacc 240

actgccgcac acgactcctg aacc atg ggg gaa aac gag gat gag aag cag 291

Met Gly Glu Asn Glu Asp Glu Lys Gln

| gcc | cag | gcg | ggg | cag | gtt | ttt | gag | aac | ttt | gtc | cag | gca | tcc | acg | tgc | 339 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Gln | Ala | Gly | Gln | Val | Phe | Glu | Asn | Phe | Val | Gln | Ala | Ser | Thr | Cys | |
| 10 | | | | | 15 | | | | | 20 | | | | | 25 | |
| | | | | | | | | | | | | | | | | |
| aaa | ggt | acc | ctc | cag | gcc | ttc | aac | att | ctc | aca | cga | cac | ctg | gac | cta | 387 |
| Lys | Gly | Thr | Leu | Gln | Ala | Phe | Asn | Ile | Leu | Thr | Arg | His | Leu | Asp | Leu | |
| | , | | | 30 | | | | | 35 | | | | | 40 | | |
| | | | | | | | | | | | | | | | | |
| gac | cct | ctg | gac | cac | aga | aac | ttt | tat | tcc | aag | ctc | aag | tcc | aag | gtg | 435 |
| Asp | Pro | Leu | Asp | His | Arg | Asn | Phe | Tyr | Ser | Lys | Leu | Lys | Ser | Lys | Val | |
| | | | 45 | | | | | 50 | | | | | 55 | • | | |
| | | | | | | | | ٠ | | | | | | | | |
| acc | acc | tgg | aaa | gcc | aaa | gcc | ctg | tgg | tac | aaa | ttg | gat | aag | cgt | ggt | 483 |
| Thr | Thr | Trp | Lys | Ala | Lys | Ala | Leu | Trp | Tyr | Lys | Leu | Asp | Lys | Arg | Gly | |
| | | 60 | | • | | | 65 | | | | | 70 | | | | |
| | | | | | | | | | | | | | | | | |
| tcc | cac | aaa | gag | tat | aag | cga | ggg | aag | tcg | tgc | acg | aac | acc | aag | tgt | 531 |
| Ser | His | Lys | Glu | Tyr | Lys | Arg | G1y | Lys | Ser | Cys | Thr | Asn | Thr | Lys | Cys | |
| | 75 | ٠ | ÷ | | | 80 | | | | | 85 | • | | | | |
| | | | | | | | | | | | | | | | | |
| ctc | ata | gtt | ggg | gga | gga | ccc | tgt | ggc | ttg | cgc | act | gcc | att | gaa | ctt | 579 |
| Leu | Ile | Val | Gly | Gly | Gly | Pro | Cys | Gly | Leu | Arg | Thr | Ala | Ile | Glu | Leu | |
| 90 | | | | | 95 | | | | | 100 | | | | | 105 | |
| | | | | | | | | | | | | | | | | |

gcc tac ctg gga gcc aaa gtg gtc gtg gtg gag aag agg gac tcc ttc 627

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Ala Tyr Leu Gly Ala Lys Val Val Val Glu Lys Arg Asp Ser Phe
110 115 120

tcc cgg aac aac gtg cta cac ctc tgg cct ttc acc atc cat gac ctt

Ser Arg Asn Asn Val Leu His Leu Trp Pro Phe Thr Ile His Asp Leu

125

130

135

cgg ggc ctg gga gcc aag aag ttc tat ggg aag ttc tgt gct ggc tcc

723

Arg Gly Leu Gly Ala Lys Lys Phe Tyr Gly Lys Phe Cys Ala Gly Ser

140

145

150

atc gac cat atc agt att cgc caa cta cag ctc atc cta ttc aag gtg

771

Ile Asp His Ile Ser Ile Arg Gln Leu Gln Leu Ile Leu Phe Lys Val

155

160

165

gcc ctg atg ctg gga gtt gaa atc cat gtg aat gtg gag ttc gtg aag 819
Ala Leu Met Leu Gly Val Glu Ile His Val Asn Val Glu Phe Val Lys
170 175 180 185

gtt cta gag cct cct gaa gat caa gaa aat caa aaa att ggc tgg cgg

Val Leu Glu Pro Pro Glu Asp Gln Glu Asn Gln Lys Ile Gly Trp Arg

190 195 200

gca gaa ttt ctc cct aca gac cat tct ctg tcg gag ttt gag ttt gac 915
Ala Glu Phe Leu Pro Thr Asp His Ser Leu Ser Glu Phe Glu Phe Asp
205 210 215

1251

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| gtc | atc | att | ggt | gcc | gat | ggc | cgc | agg | aac | acc | ctg | gaa | ggg | ttc | aga | 963 |
|-----|------|-------|-----|-----|------|------|-----|-------|-----|------|--------|-----|-----|-------|-------|------|
| Val | Ile | Ile | Gly | Ala | Asp | Gly | Arg | Arg | Asn | Thr | Leu | Glu | Gly | Phe | Arg | |
| | | 220 | | | | | 225 | | | | | 230 | | | | |
| | | • | | | , | | | | | | | | | | | |
| aga | aaa | gaa | ttc | cgt | ggg | aag | ctg | gcg | att | gcc | atc | acc | gcc | aac | ttc | 1011 |
| Arg | Lys | Glu | Phe | Arg | Gly | Lys | Leu | Ala | Ile | Ala | Ile | Thr | Ala | Asn | Phe | |
| | 235 | | | `- | | 240 | | | | | 245 | | | | | • |
| | | | | | | | | | | | | | | | | |
| | | | aac | | | | | | | | | | | | | 1059 |
| | Asn | Arg | Asn | Ser | | Ala | Glu | Ala | Lys | | Glu | Glu | Ile | Ser | | ٠. |
| 250 | | | | | 255 | | | | | 260 | • | | | | 265 | |
| ~+~ | ant. | ++0 | atc | ++0 | oot | 000 | 000 | +++ | +++ | 000 | go o i | att | 999 | a a a | g a a | 1107 |
| | - | | Ile | · | | | | | | | | | | | | 1101 |
| Val | піа | 1 116 | 116 | 270 | VSII | GIII | гуз | 1 116 | 275 | GIII | nsp | Deu | Lys | 280 | | |
| | | | | 210 | | | | | 210 | | | | | 200 | | • |
| aca | ggc | ata | gat | ctt | gag | aac | att | gtt | tac | tac | aag | gac | tgc | acc | cac | 1155 |
| | | | Asp | | | | | | | | | , | | | | |
| | | | 285 | | | | | 290 | | | | | 295 | | | |
| | | | | | | | | | | | | | | | • | |
| tat | ttt | gta | atg | aca | gcc | aag | aag | cag | agc | ctg | ctc | gac | aaa | ggt | gtc | 1203 |
| Tyr | Phe | Val | Met | Thr | Ala | Lys | Lys | Gln | Ser | Leu | Leu | Asp | Lys | Gly | Val | |
| | | 300 | | | | | 305 | | | | | 310 | | | | |
| | | | | | | | | | | | | | | | | |

atc att aac gac tac atc gac aca gag atg ctg ctg tgt gcg gag aac

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Ile Ile Asn Asp Tyr Ile Asp Thr Glu Met Leu Leu Cys Ala Glu Asn
315 320 325

gtg aac caa gac aac ctg cta tcc tat gcc cgg gaa gct gca gac ttt 1299
Val Asn Gln Asp Asn Leu Leu Ser Tyr Ala Arg Glu Ala Ala Asp Phe
330 335 340 345

gcc acc aac tac cag ctg cca tcc tta gac ttt gcc atg aac cac tat

1347

Ala Thr Asn Tyr Gln Leu Pro Ser Leu Asp Phe Ala Met Asn His Tyr

. 350

355

360

ggg cag cct gat gtg gcc atg ttt gac ttt acc tgc atg tat gcc tca 1395

Gly Gln Pro Asp Val Ala Met Phe Asp Phe Thr Cys Met Tyr Ala Ser

365 370 375

gag aac gcg gcc ctg gtg cgg gag cgg cag gcg cac cag ctg ctc gtg 1443
Glu Asn Ala Ala Leu Val Arg Glu Arg Gln Ala His Gln Leu Leu Val
380 385 390

gcc ctt gtg ggt gac agc ttg ctt gag cca ttt tgg ccc atg ggt aca

1491

Ala Leu Val Gly Asp Ser Leu Leu Glu Pro Phe Trp Pro Met Gly Thr

395

400

405

ggc tgt gcc cgt ggc ttc ctg gca gcc ttt gac acg gca tgg atg gtg 1539

Gly Cys Ala Arg Gly Phe Leu Ala Ala Phe Asp Thr Ala Trp Met Val

410 415 420 425

| aag | agc | tgg | aac | cag | ggc | acc | cct | ссс | ctg | gag | ctg | ctg | gct | gaa | agg | 1587 |
|-------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-------|-----|------|------|
| Lys | Ser | Trp | Asn | Gln | Gly | Thr | Pro | Pro | Leu | Glu | Leu | Leu | Ala | Glu | Arg | |
| | | | | 430 | | | | | 435 | | | | | 440 | | |
| | | | | | | | | | | | | | | | | |
| gaa | agt | ctc | tac | cgg | ctg | tta | cct | cag | aca | acc | ccg | gag | aac | atc | aac | 1635 |
| Glu | Ser | Leu | Tyr | Arg | Leu | Leu | Pro | Gln | Thr | Thr | Pro | Glu | Asn | Ile | Asn | |
| | | | 445 | | | | | 450 | | | | | 455 | | • | |
| | | | | | | | | | | | | | | | | |
| aag | aac | ttt | gag | cag | tac | acg | ttg | gac | cca | ggg | aca | cgg | tac | cca | aac | 1683 |
| | | | Glu | | | | | | | | | | | _ | | |
| _, | | 460 | | | • | | 465 | _ | | | | 470 | | | | |
| | | 100 | | | | | | | | | | | | | | |
| . 4 . | | 4 | | ++ | | | | | | ~+~ | 000 | aat | . ++~ | tot | at a | 1731 |
| | | | cac | | | | | | | | | | | | | 1101 |
| Leu | Asn | Ser | His | Суѕ | Val | | Pro | His | GIn | Val | | | Leu | Tyr | lie | |
| | 475 | | | | | 480 | | | | | 485 | • | | | | |
| | | : . | | | | | | | | | | | | | | |
| act | aag | gag | ctg | gag | cac | tac | cct | ctc | gag | aga | ctg | ggc | tcg | gtg | agg | 1779 |
| Thr | Lys | Glu | Leu | Glu | His | Tyr | Pro | Leu | Glu | Arg | Leu | Gly | Ser | Val | Arg | |
| 490 | | | | , | 495 | | | | | 500 | | | | | 505 | |
| | | | | | | | | | | | | | | | | |
| aga | tct | gtc | aac | ctc | tcc | agg | aag | gag | tca | gat | atc | cgg | ccc | agc | aag | 1827 |
| Arg | Ser | Val | Asn | Leu | Ser | Arg | Lys | Glu | Ser | Asp | Ile | Arg | Pro | Ser | Lys | |
| | | | | 510 | | | | | 515 | | | | | 520 | | |
| | | | | | | | | | | | | | | | / | |
| | | | | | | | | | | | | | | | | |

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Leu Leu Thr Trp Cys Gln Gln Gln Thr Glu Gly Tyr Gln His Val Asn
525 530 535

gtc acc gac ctg acc aca tcc tgg cgc agt ggg ttg gcc ctg tgt gcc 1923

Val Thr Asp Leu Thr Thr Ser Trp Arg Ser Gly Leu Ala Leu Cys Ala

540

545

550

atc atc cac cgc ttc cgg cct gag ctc atc aac ttt gac tct ttg aat 1971

Ile Ile His Arg Phe Arg Pro Glu Leu Ile Asn Phe Asp Ser Leu Asn

555 560 565

gaa gat gat gct gtg gag aac aac cag ctc gca ttt gat gtg gcc gag 2019 Glu Asp Asp Ala Val Glu Asn Asn Gln Leu Ala Phe Asp Val Ala Glu 570 575 580 585

cga gag ttt ggg atc cct cca gtg acc acg ggc aaa gag atg gca tct 2067

Arg Glu Phe Gly Ile Pro Pro Val Thr Thr Gly Lys Glu Met Ala Ser

590 595 600

gcc cag gag cct gac aag ctc agc atg gtc atg tac ctc tcc aag ttc 2115

Ala Gln Glu Pro Asp Lys Leu Ser Met Val Met Tyr Leu Ser Lys Phe
605 610 615

tac gag ctc ttc cgg ggc acc cca ctg agg ccc gtg gat tct tgg cgc 2163

Tyr Glu Leu Phe Arg Gly Thr Pro Leu Arg Pro Val Asp Ser Trp Arg
620 625 630

2499

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| aa | a a | ac | tat | gga | gaa | aat | gct | gac | ctc | agc | ttg | gcc | aaa | tca | tcc | att | 2211 |
|----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|------|
| Ly | s A | sn | Tyr | Gly | Glu | Asn | Ala | Asp | Leu | Ser | Leu | Ala | Lys | Ser | Ser | Ile | |
| | 6 | 35 | | | | | 640 | | | | • | 645 | | | | | |
| | | | | | | | | | | | | • | | | | | |
| tc | t a | at | aac | tat | ctc | aac | ctc | aca | ttt | cca | agg | aag | agg | act | cca | cgg | 2259 |
| Se | r.A | lsn | Asn | Tyr | Leu | Asn | Leu | Thr | Phe | Pro | Arg | Lys | Arg | Thr | Pro | Arg | |
| 65 | 0 | | | | | 655 | | | | | 660 | | | | | 665 | |
| | | | | | | | | | | | ٠ | | | | | | |
| gt | g g | gat | ggt | caa | acc | gga | gag | aat | gac | atg | aac | aaa | cgg | aga | cgg | aag | 2307 |
| Va | 1 A | lsp | Gly | Gln | Thr | Gly | Glu | Asn | Asp | Met | Asn | Lys | Arg | Arg | Arg | Lys | |
| | ٠ | | | | 670 | | | | | 675 | | | | | 680 | • | |
| | | | | | | | | | | | | | | | | | |
| gg | c t | ttc | acc | aac | ctg | gac | gag | cct | tca | aac | ttt | tcc | agc | cgt | agc | ttg | 2355 |
| G1 | y F | Phe | Thr | Asn | Leu | Asp | Glu | Pro | Ser | Asn | Phe | Ser | Ser | Arg | Ser | Leu | |
| | | | | 685 | | | | | 690 | | | | | 695 | | | |
| | | | | | | | | | | | | | | | | | |
| gg | c t | tcc | aat | caa | gag | tgt | ggg | agc | agt | aag | gaa | ggt | gga | aat | cag | aac. | 2403 |
| G1 | y S | Ser | Asn | Gln | Glu | Cys | G1y | Ser | Ser | Lys | Glu | Gly | Gly | Asn | Gln | Asn | |
| | | | 700 | | | | | 705 | | | | | 710 | | | | |
| | | | | | | | | ٠ | | | | | | | | | |
| aa | a g | gtc | aag | tcc | atg | gcg | aat | cag | ctg | ctg | gcc | aag | ttt | gag | gag | agc | 2451 |
| Ly | s l | Val | Lys | Ser | Met | Ala | Asn | Gln | Leu | Leu | Ala | Lys | Phe | Glu | Glu | Ser | |
| | 7 | 715 | | | | | 720 | | | | | 725 | | | | | |
| | | | | | | | | | | | | | | | | | |

act cgg aac ccc tca ctc atg aag cag gaa tct atg cga aag tca ttt

| Thr | Arg | Asn | Pro | Ser | Leu | Met | Lys | Gln | Glu | Ser | Met | Arg | Lys | Ser | Phe | |
|------|------|-----|------|------|-----|-----|------|-----------|-----|-----|-----|-----|------|-----|----------|------|
| 730 | | | | | 735 | | | | | 740 | | | | | 745 | |
| | | | | | | | | | | | | | | | | |
| ccc | ctt | aac | ctg | gga | ggc | agc | gac | acg | tgt | tac | ttc | tgt | aag | aaa | cgt | 2547 |
| Pro | Leu | Asn | Leu | Gly | Gly | Ser | Asp | Thr | Cys | Tyr | Phe | Cys | Lys | Lys | Arg | |
| | | | | 750 | | | | | 755 | | | | | 760 | | |
| | | | | | | | | | | | | | | | | |
| gtg | tac | gtg | atg | gaa | cgg | ctg | agc | gcc | gag | ggc | cac | ttc | ttc | cac | cgg | 2595 |
| | | | Met | | | | | | | | | | | • | | |
| | | | 765 | | | | | 770 | | | | | 775 | | 8 | |
| | | | | | | | | ••• | | | | | | | | |
| ~~ ~ | + a+ | ++0 | 0.00 | + 00 | 200 | oto | + a+ | ~~ | | 000 | ++~ | 000 | at a | ~~~ | ~ | 2642 |
| | | | cgc | | | | | | | | | | | - | | 2643 |
| GIU | cys | | Arg | Cys | Ser | 116 | | Ата | Inr | inr | Leu | | Leu | ATA | Ala | |
| | | 780 | | | | | 785 | | | | | 790 | | | | |
| | | | | | | | | | | | | | | | , | |
| tac | acc | ttt | gac | tgc | gat | gaa | ggc | aaa | ttt | tac | tgc | aag | cct | cac | ttc | 2691 |
| Tyr | Thr | Phe | Asp | Cys | Asp | Glu | Gly | Lys | Phe | Tyr | Cys | Lys | Pro | His | Phe | |
| | 795 | | | | | 800 | | | | | 805 | | · | | | |
| • | ٠ | | | | | | | | | | | | | | | |
| att | cac | tgt | aaa | acc | aat | agc | aaa | caa | cgg | aag | aga | cgg | gca | gag | ttg | 2739 |
| Ile | His | Cys | Lys | Thr | Asn | Ser | Lys | Gln | Arg | Lys | Arg | Arg | Ala | Glu | Leu | |
| 810 | | | | | 815 | | | | | 820 | | | | | 825 | |
| | | | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | | , | 0505 |

aag caa caa aga gag gag gag gca aca tgg caa gag cag gaa gcc cct 2787

Lys Gln Gln Arg Glu Glu Glu Ala Thr Trp Gln Glu Gln Glu Ala Pro

830 835 840

3123

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| cgg | aga | gac | act | ccc | acc | gaa | agt | tct | tgc | gca | gtg | gcc | gcc | att | ggc | 2835 |
|-----|-----|-----|-------|------|------|-------|-----|-----|-------|-----|-----|-----|-----|------|-----|------|
| Arg | Arg | Asp | Thr | Pro | Thr | Glu | Ser | Ser | Cys | Ala | Val | Ala | Ala | Ile | Gly | |
| | | | 845 | | | | | 850 | | | | | 855 | | | |
| | | | | | | | | | | | | | | | | |
| acc | ctg | gaa | ggc | agc | ccc | cca | ggt | atc | tcc | acc | tcc | ttc | ttt | agg | aag | 2883 |
| Thr | Leu | Glu | Gly | Ser | Pro | Pro | Gly | Ile | Ser | Thr | Ser | Phe | Phe | Arg | Lys | |
| | | 860 | | | | | 865 | | | | | 870 | | | | |
| | | | | | | | | | | | | | | | | |
| gtg | ctg | ggc | tgg | ccc | ctc | agg | ctg | ccg | agg | gac | ctg | tgt | aac | tgg | atg | 2931 |
| Val | Leu | Gly | Trp | Pro | Leu | Arg | Leu | Pro | Arg | Asp | Leu | Cys | Asn | Trp | Met | |
| | 875 | | | | | 880 | | | | | 885 | | | 1 | | |
| | | | | | | | | | | | | | | | | |
| cag | gga | ctc | ctg | caa | gct | gct | ggc | ctc | cat | atc | agg | gac | aat | gct | tac | 2979 |
| Gln | Gly | Leu | Leu | Gln | Ala | Ala | Gly | Leu | His | Ile | Arg | Asp | Asn | Ala | Tyr | |
| 890 | | | | | 895 | | | | | 900 | | | | | 905 | |
| | | | | | | | | | | | | | | | | |
| aac | tac | tgc | tac | atg | tac | gag | ctc | ctg | agc | ctg | ggg | ctg | cca | ctc | ctc | 3027 |
| | | | | Met | | | | | | | | | | | | |
| | -,- | -,- | -,- | 910 | -,- | | | | 915 | | | | | 920 | | |
| , | | | | 010 | | | | | 010 | | | | | 0.20 | | |
| +~~ | ~~~ | ++0 | + 0 + | ~~ ~ | at o | o t a | avo | aoo | a t a | too | ann | raa | tot | as a | aac | 3075 |
| | | | | gag | | | | | | | | | | | | 3013 |
| ırp | Ala | rne | | Glu | vai | Leu | нта | | Met | ıyr | Arg | GIU | | GTŊ | ату | |
| | | | 925 | | | | | 930 | | | | | 935 | | | |
| | | | • | | | | | | | | | | | | | |

tcc ctc gag agc atc tgc aac tgg gtg ctc agg tgc ttc cca gtc aag

Ser Leu Glu Ser Ile Cys Asn Trp Val Leu Arg Cys Phe Pro Val Lys 940 945 950

| ctc | cgc | tga | catggctggc | tgccccaaag | tgccttcaca | tttccaggga | | 3172 |
|------|------------|------|------------|------------|------------|------------|------------|---------------|
| Leu | Arg 955 | , | | | | | | |
| ggc | ttca | gat | ggcagtgcgt | ttgcagtttg | ctcaggctct | ggccaggaag | cctagcattc | 3232 |
| tcta | aagc | aat | tagctcaaag | ccaaagaatt | tcacatgggc | cacctccgcc | tggccttatc | 3292 |
| agg | gtga | aca | tctactcacg | gtgctagggc | cagggatgat | atgaaggatc | ttttctatag | 3352 |
| ctt | tgtg | agc | catacttctg | ggtttacatt | tcaatttttt | taattttaat | tagcccagag | 3412 |
| aaa | gcat | ttt | tttctatgag | tgtcaatttt | tctaaacatg | ggtttgaagc | ttataaccag | 3 <u>4</u> 72 |
| ttt | tata | aac | cccttgaaca | ctgcagtgag | ttatcaaagc | cactgcctgc | aaagtggatg | 3532 |
| att | taag | att | ttacacgcat | gaaaatgagt | gtgccatctc | ctgaccagtg | ccttttgact | 3592 |
| tag | gtac | cca | gatgccactt | gtcagcagca | ggatactttt | tacaacacga | aagcataatt | 3652 |
| att | ttag | gaag | aagagagtag | aagggcagaa | tagaattcaa | cttacagaag | cacggagcag | 3712 |
| tgt | gtgg | gttg | gctgttatct | gtcccctgg | gaggaggact | gttttgctcc | cttgttttga | 3772 |

| tgttaaacag | tagcttaaag | gctttcccc | ccataccaac | tcacagccaa | atgacaaaga | 3832 |
|------------|------------|------------|-----------------|------------|------------|------|
| accgtggggt | ttcaacagat | tctacaaaca | tgcattttcc | cttcccacta | atgggcactg | 3892 |
| cagggaaagc | ccattggcat | ttgaccatgg | agctgatgca | gtgccaaaga | tgagetettt | 3952 |
| caactgatgg | cattttagcc | cctgtggctc | ccagcggatc | ccccagcccg | ggctgcaggc | 4012 |
| tgagccaagg | ctgtgcaggg | tccatattgg | tcaggccaag | tggagtggaa | gactctgtcc | 4072 |
| acttatgtgg | tgtcctttgg | gactgagggg | gtttgttagc | acatcagget | attgctggga | 4132 |
| agcgtggcct | gcccagtgag | cattgcctgt | , ggacatcctg | actgcttagc | tgctccgctg | 4192 |
| ccacacatat | gtggtcaaaa | cagaaaccaa | tttcacactg | ccctgggaaa | ggaatgggtc | 4252 |
| tgacctccag | gggaagctct | accatatctt | gactggcagg | gaaggctggg | agtggaagct | 4312 |
| atttatggac | tgatccaaag | gacatatgca | tgagtaaggg | taaaaatgag | catgcaggtc | 4372 |
| cacctgtgtt | cttactctgg | gtatctagaa | gagtcctcag | ctctccctac | tccacgctgc | 4432 |
| ctagacatac | acagctgcag | ggtctggctg | aacaatcaag | gggccgccag | agaaaggcca | 4492 |
| tctacggtgc | gcagtgtatc | tggagttgct | gggcccaaga | tagctctgtg | gagttatcac | 4552 |

3 9 / 5 9

| tagagatgcc | tctggattaa | ctaagaggtg | tgcctgggtg | tgggtgagga | gtcagaacct | 4612 |
|------------|------------|------------|------------|------------|------------|------|
| ttgagagctt | tgagatgaca | gtttctatgg | ggcgggaaga | aggaggtgca | tttctacaaa | 4672 |
| cacttccctg | aaatccttgg | gaaaaacaga | ggcatggccg | tggccaactc | tgtgggaact | 4732 |
| ggcgcctctg | tccttgttgg | cactgttctc | agtccgatga | cttgcattgt | gttttctcca | 4792 |
| atttttgctg | ggattttaat | gttcagcatg | gtgggaggaa | cccttgattc | cttttgtttg | 4852 |
| agtatagaaa | gtaaattttt | gaggtcatga | tgtgaacggc | catgttattg | tgattatctt | 4912 |
| cagctcagga | taggctgaga | tgctttgtgg | agtgttccat | gaagcccgag | toggaatoto | 4972 |
| tgactgtcgt | gtacagccat | aaggagactg | gtttgaatta | ctgtggcgag | acagggcgtg | 5032 |
| cctgtcagaa | atctgagatg | tttgtacgct | ctgagatgtt | gaacetttet | ggtgggcagc | 5092 |
| accgacaccc | aggggtggac | ccccgaggat | gaatgcctct | aggcctccgc | aacatattca | 5152 |
| agaatgaatg | ggagacgcta | gagtaaaatg | ggggcagaga | ggatatcagg | gagcaagatg | 5212 |
| caaactgtgt | gcatccactc | tcgtaaacaa | gtagctggtc | acaaccagaa | aggttcatct | 5272 |
| ctcctaagca | aacagcgact | ctttcagagg | aagtttccct | ctttcaatcg | tggccttatt | 5332 |

| ttcaactccg | gtgccttctc | gtgatgttaa | tcatttcctt | ttttccccac | actaagetet | 5392 |
|------------|------------|------------|------------|------------|------------|------|
| cttttctatc | tttctctctc | tttccaatct | tacgccatgg | ccatcagttc | atttcagcct | 5452 |
| tccagtgcta | cacccacttc | ttggctgaca | cacttctgct | ctaaggtgac | tggttttctt | 5512 |
| gccaattttc | aaagagtggt | actaaccccc | aacccgcttt | ccgcaccccg | tcctctccgc | 5572 |
| cagcagtact | ggttgcacta | actgtgagtg | tcttgcatac | tgatggactc | atttggtggc | 5632 |
| atggttggct | aacagcatgg | cggggggtgt | tcagcttgag | acccatgcct | gtgttcattt | 5692 |
| cccatggagc | tggcagcctg | gtctacccca | agtgcatgcc | ccgcctctcc | tetetecett | 5752 |
| gggtctgcct | gcgtgcatgc | ttctccagtt | gcgtctgcga | agctacctac | tttcttggga | 5812 |
| gggtcgacct | tgatcatgaa | acaataccat | gagggggcct | ctgtcacctt | tgaaaagaac | 5872 |
| actttttgag | cagcctcaaa | aagctcatac | ataccagcgc | cttcttaaat | tggctctaat | 5932 |
| gtaaagattg | ttaatgtcat | ttatcaaaac | cataggtgat | tatttggagg | gatttaaaaa | 5992 |
| acttaattac | tctcaggcct | catcccaagc | ttgacacatg | ctctgtaggt | tgaacacata | 6052 |
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6172 ttgctgtggc tcttcttatc tcccttggct cataaagccc cagatgatgc cagagcttca 6232 attagageca teateateee aggeagggat atetttgaga aatgaeteag tteageecea ggcccctgtg actctgctta aagcacacat ttctgctgac tcttgtacct ggggcagcag 6292 6352 gataatcacc aacacactet taacgagaaa caacacacca agcaccgtgg agctgtccta 6412 ggcaacactc gcggtctcag gctgcggtgg gcgtctgtcc tgcatgtggc ccagaccacc ctgaccccg ggcctgcctg cctggccctg catgctgcac gctcactgta tttgtgcaga 6472 6532 tectggecag tacaaagteg ttgetettgt ettatettet ettacagagt eteceteeet ttatagaatg tcaaccaaag agtgccctcc tcccctctca gcctcctctt tagctagcct 6592 6652 ccccatctca tcacaacgca tgtctgtgac ctttggtaat catttacagt gccacacgga 6712 accetgtatt ttgcacacag caaaacaaac aatgtttage tttatttatg gtatttgatg 6742 ctgtaaatgg aaataaatat tgttctttat

<210> 6

<211> 955

30

42/59

<212> PRT

<213> Homo sapiens

<400> 6

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5 1 10 15

Glu Asn Phe Val Gln Ala Ser Thr Cys Lys Gly Thr Leu Gln Ala Phe 20 25

Asn Ile Leu Thr Arg His Leu Asp Leu Asp Pro Leu Asp His Arg Asn 35 40 45

Phe Tyr Ser Lys Leu Lys Ser Lys Val Thr Thr Trp Lys Ala Lys Ala 50 55 60

Leu Trp Tyr Lys Leu Asp Lys Arg Gly Ser His Lys Glu Tyr Lys Arg 65 70 75 80

Gly Lys Ser Cys Thr Asn Thr Lys Cys Leu Ile Val Gly Gly Gly Pro 85 90 95

Cys Gly Leu Arg Thr Ala Ile Glu Leu Ala Tyr Leu Gly Ala Lys Val 100 105 110

Val Val Val Glu Lys Arg Asp Ser Phe Ser Arg Asn Asn Val Leu His

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115 120 125

Leu Trp Pro Phe Thr Ile His Asp Leu Arg Gly Leu Gly Ala Lys Lys

130 135 140

Phe Tyr Gly Lys Phe Cys Ala Gly Ser Ile Asp His Ile Ser Ile Arg
145 150 155 160

Gln Leu Gln Leu Ile Leu Phe Lys Val Ala Leu Met Leu Gly Val Glu 165 170 175

Ile His Val Asn Val Glu Phe Val Lys Val Leu Glu Pro Pro Glu Asp 180 185 190

Gln Glu Asn Gln Lys Ile Gly Trp Arg Ala Glu Phe Leu Pro Thr Asp

195 200 205

His Ser Leu Ser Glu Phe Glu Phe Asp Val Ile Ile Gly Ala Asp Gly
210 215 220

Arg Arg Asn Thr Leu Glu Gly Phe Arg Arg Lys Glu Phe Arg Gly Lys
225 230 235 240

Leu Ala Ile Ala Ile Thr Ala Asn Phe Ile Asn Arg Asn Ser Thr Ala
245 250 255

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Glu Ala Lys Val Glu Glu Ile Ser Gly Val Ala Phe Ile Phe Asn Gln
260 265 270

Lys Phe Phe Gln Asp Leu Lys Glu Glu Thr Gly Ile Asp Leu Glu Asn
275
280
-285

Ile Val Tyr Tyr Lys Asp Cys Thr His Tyr Phe Val Met Thr Ala Lys
290 295 300

Lys Gln Ser Leu Leu Asp Lys Gly Val Ile Ile Asn Asp Tyr Ile Asp 305 310 315 320

Thr Glu Met Leu Cys Ala Glu Asn Val Asn Gln Asp Asn Leu Leu
325 330 335

Ser Tyr Ala Arg Glu Ala Ala Asp Phe Ala Thr Asn Tyr Gln Leu Pro 340 345 350

Ser Leu Asp Phe Ala Met Asn His Tyr Gly Gln Pro Asp Val Ala Met 355 360 365

Phe Asp Phe Thr Cys Met Tyr Ala Ser Glu Asn Ala Ala Leu Val Arg 370 375 380

Glu Arg Gln Ala His Gln Leu Leu Val Ala Leu Val Gly Asp Ser Leu 385 390 395 400

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Leu Glu Pro Phe Trp Pro Met Gly Thr Gly Cys Ala Arg Gly Phe Leu
405 410 415

Ala Ala Phe Asp Thr Ala Trp Met Val Lys Ser Trp Asn Gln Gly Thr
420 425 430

Pro Pro Leu Glu Leu Leu Ala Glu Arg Glu Ser Leu Tyr Arg Leu Leu
435 440 445

Pro Gln Thr Thr Pro Glu Asn Ile Asn Lys Asn Phe Glu Gln Tyr Thr
450 455 460

Leu Asp Pro Gly Thr Arg Tyr Pro Asn Leu Asn Ser His Cys Val Arg
465 470 475 480

Pro His Gln Val Lys His Leu Tyr Ile Thr Lys Glu Leu Glu His Tyr
485 490 495

Pro Leu Glu Arg Leu Gly Ser Val Arg Arg Ser Val Asn Leu Ser Arg
500 505 510

Lys Glu Ser Asp Ile Arg Pro Ser Lys Leu Leu Thr Trp Cys Gln Gln
515 520 525

Gln Thr Glu Gly Tyr Gln His Val Asn Val Thr Asp Leu Thr Thr Ser

530

535

540

Trp Arg Ser Gly Leu Ala Leu Cys Ala Ile Ile His Arg Phe Arg Pro 545 550 555 560

Glu Leu Ile Asn Phe Asp Ser Leu Asn Glu Asp Asp Ala Val Glu Asn
565 570 575

Asn Gln Leu Ala Phe Asp Val Ala Glu Arg Glu Phe Gly Ile Pro Pro
580 585 590

Val Thr Thr Gly Lys Glu Met Ala Ser Ala Gln Glu Pro Asp Lys Leu
595 600 605

Ser Met Val Met Tyr Leu Ser Lys Phe Tyr Glu Leu Phe Arg Gly Thr
610 620

Pro Leu Arg Pro Val Asp Ser Trp Arg Lys Asn Tyr Gly Glu Asn Ala 625 630 635 640

Asp Leu Ser Leu Ala Lys Ser Ser Ile Ser Asn Asn Tyr Leu Asn Leu 645 650 655

Thr Phe Pro Arg Lys Arg Thr Pro Arg Val Asp Gly Gln Thr Gly Glu
660 665 670

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Asn Asp Met Asn Lys Arg Arg Lys Gly Phe Thr Asn Leu Asp Glu 675 680 685

Pro Ser Asn Phe Ser Ser Arg Ser Leu Gly Ser Asn Gln Glu Cys Gly
690 695 700

Ser Ser Lys Glu Gly Gly Asn Gln Asn Lys Val Lys Ser Met Ala Asn 705 710 715 720

Gln Leu Leu Ala Lys Phe Glu Glu Ser Thr Arg Asn Pro Ser Leu Met
725 730 735

Lys Gln Glu Ser Met Arg Lys Ser Phe Pro Leu Asn Leu Gly Gly Ser
740 745 750

Asp Thr Cys Tyr Phe Cys Lys Lys Arg Val Tyr Val Met Glu Arg Leu
755 760 765

Ser Ala Glu Gly His Phe Phe His Arg Glu Cys Phe Arg Cys Ser Ile
770 780

Cys Ala Thr Thr Leu Arg Leu Ala Ala Tyr Thr Phe Asp Cys Asp Glu
785 790 795 800

Gly Lys Phe Tyr Cys Lys Pro His Phe Ile His Cys Lys Thr Asn Ser 805 810 815

Lys Gln Arg Lys Arg Arg Ala Glu Leu Lys Gln Gln Arg Glu Glu Glu 820 825 830

Ala Thr Trp Gln Glu Gln Glu Ala Pro Arg Arg Asp Thr Pro Thr Glu 835 840 845

Ser Ser Cys Ala Val Ala Ala Ile Gly Thr Leu Glu Gly Ser Pro Pro 850 855 860

Gly Ile Ser Thr Ser Phe Phe Arg Lys Val Leu Gly Trp Pro Leu Arg 865 870 875 880

Leu Pro Arg Asp Leu Cys Asn Trp Met Gln Gly Leu Leu Gln Ala Ala 885 890 895

Gly Leu His Ile Arg Asp Asn Ala Tyr Asn Tyr Cys Tyr Met Tyr Glu 900 905 910

Leu Leu Ser Leu Gly Leu Pro Leu Leu Trp Ala Phe Ser Glu Val Leu
915 920 925

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51/59

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